

AMENDMENTS TO THE CLAIMS

This listing of the claims will replace all prior listings and versions of claims in the application:

- 1-52. (cancelled)
53. (previously presented) A method for treating nucleic acid, comprising:
- a) providing:
 - i) a purified archael FEN-1 endonuclease; and
 - ii) a nucleic acid substrate;
 - b) treating said nucleic acid substrate under conditions such that said substrate forms one or more cleavage structures; and
 - c) reacting said endonuclease with said cleavage structures so that one or more cleavage products are produced.
54. (previously presented) The method of Claim 53, wherein said archael FEN-1 endonuclease comprises an *Archaeoglobus fulgidus* FEN-1 endonuclease.
55. (previously presented) The method of Claim 54, wherein said *Archaeoglobus fulgidus* FEN-1 endonuclease comprises SEQ ID NO:179.
56. (previously presented) The method of Claim 53, wherein said purified archael FEN-1 endonuclease is provided as part of a mixture, said mixture further comprising a second structure-specific nuclease, wherein said reacting said endonuclease with said cleavage structures in step c) comprises reacting said cleavage structure with said mixture.
57. (previously presented) The method of Claim 56, wherein said second structure-specific nuclease comprises a polymerase.
58. (previously presented) The method of Claim 57, wherein said polymerase comprises a thermostable polymerase.
59. (previously presented) The method of Claim 58, wherein said thermostable polymerase comprises a thermostable DNA polymerase altered in amino acid sequence such that it exhibits reduced DNA synthetic activity from that of a corresponding wild-type DNA polymerase but retains substantially the same 5' nuclease activity of said wild-type DNA polymerase.

60. (previously presented) The method of Claim 53, wherein said nucleic acid of step (a) is substantially single stranded.
61. (previously presented) The method of Claim 53, wherein said nucleic acid of step (a) is double stranded.
62. (previously presented) The method of Claim 61, wherein said treating of step (b) comprises:
- i) rendering said double-stranded nucleic acid substantially single-stranded; and
 - ii) exposing said single-stranded nucleic acid to conditions such that said single-stranded nucleic acid has secondary structure.
63. (previously presented) The method of Claim 62, wherein said double stranded nucleic acid is rendered substantially single-stranded by the use of increased temperature.
64. (previously presented) The method of Claim 53, further comprising the step of detecting at least one of said one or more cleavage products.
65. (previously presented) A method for treating nucleic acid, comprising:
- a) providing:
 - i) a purified FEN-1 endonuclease selected from the group consisting of *Pyrococcus woesei* FEN-1 endonuclease, *Methanococcus jannaschii* FEN-1 endonuclease, *Pyrococcus furiosus* FEN-1 endonuclease, *Methanobacterium thermoautotrophicum* FEN-1 endonuclease; and chimerical FEN-1 endonuclease; and
 - ii) a nucleic acid substrate;
 - b) treating said nucleic acid substrate under conditions such that said substrate forms one or more cleavage structures; and
 - c) reacting said endonuclease with said cleavage structures so that one or more cleavage products are produced.
66. (previously presented) The method of Claim 65, wherein said nucleic acid substrate comprises DNA.

67. (previously presented) The method of Claim 65, wherein said *Methanococcus jannaschii* FEN-1 endonuclease comprises SEQ ID NO:111.
68. (previously presented) The method of Claim 65, wherein said *Methanobacterium thermoautotrophicum* FEN-1 endonuclease comprises SEQ ID NO:183.
69. (previously presented) The method of Claim 65, wherein said *Pyrococcus furiosus* FEN-1 endonuclease comprises SEQ ID NO:115.
70. (previously presented) The method of Claim 65, wherein said chimerical FEN-1 endonuclease comprises at least a portion of an archael FEN-1 endonuclease.
71. (previously presented) The method of Claim 70, wherein said archael FEN-1 endonuclease comprises a *Archaeoglobus fulgidus* FEN-1 endonuclease.
72. (previously presented) The method of Claim 65, wherein said chimerical FEN-1 endonuclease comprises at least a portion of a endonuclease selected from the group consisting of *Pyrococcus woesei* FEN-1 endonuclease, *Methanococcus jannaschii* FEN-1 endonuclease, and *Methanobacterium thermoautotrophicum* FEN-1 endonuclease.
73. (previously presented) The method of Claim 65, wherein said purified FEN-1 endonuclease is provided as part of a mixture, said mixture further comprising a second structure-specific nuclease, wherein said reacting said endonuclease with said cleavage structures in step c) comprises reacting said cleavage structure with said mixture.
74. (previously presented) The method of Claim 73, wherein said second structure-specific nuclease comprises a polymerase.
75. (previously presented) The method of Claim 74, wherein said polymerase comprises a thermostable polymerase.

76. (withdrawn) The method of Claim 75, wherein said thermostable polymerase comprises a thermostable DNA polymerase altered in amino acid sequence such that it exhibits reduced DNA synthetic activity from that of a corresponding wild-type DNA polymerase but retains substantially the same 5' nuclease activity of said wild-type DNA polymerase.

77. (previously presented) The method of Claim 65, wherein said nucleic acid of step (a) is substantially single stranded.

78. (previously presented) The method of Claim 65, wherein said nucleic acid of step (a) is double stranded.

79. (previously presented) The method of Claim 78, wherein said treating of step (b) comprises:

- i) rendering said double-stranded nucleic acid substantially single-stranded; and
- ii) exposing said single-stranded nucleic acid to conditions such that said single-stranded nucleic acid has secondary structure.

80. (previously presented) The method of Claim 79, wherein said double stranded nucleic acid is rendered substantially single-stranded by the use of increased temperature.

81. (previously presented) The method of Claim 65, further comprising the step of detecting at least one of said one or more cleavage products.

82. (previously presented) A method for treating nucleic acid, comprising:

- a) providing:
 - i) a first structure-specific nuclease comprising a purified FEN-1 endonuclease; and
 - ii) a nucleic acid substrate;
- b) treating said nucleic acid substrate with increased temperature such that said substrate is substantially single-stranded;
- c) reducing said temperature under conditions such that said single-stranded substrate forms one or more folded cleavage structures;
- d) reacting said first nuclease with said folded cleavage structures so that one or more cleavage products are produced; and

e) detecting said one or more cleavage products.

83. (previously presented) The method of Claim 82, wherein said first nuclease is provided in a solution containing manganese.

84. (previously presented) The method of Claim 82, wherein said first nuclease is selected from the group consisting *Pyrococcus woesei* FEN-1 endonuclease, *Pyrococcus furiosus* FEN-1 endonuclease, *Methanococcus jannaschii* FEN-1 endonuclease, *Methanobacterium thermoautotrophicum* FEN-1 endonuclease, *Archaeoglobus fulgidus* FEN-1 endonuclease, and chimerical FEN-1 endonucleases.

85. (previously presented) The method of Claim 82, wherein said first nuclease is provided as part of a mixture, said mixture further comprising a second structure-specific nuclease, wherein said reacting said first nuclease with said folded cleavage structures in step d) comprises reacting said folded cleavage structure with said mixture.

86. (previously presented) The method of Claim 85, wherein said second nuclease is selected from the group consisting of the Cleavase® BN enzyme, *Thermus aquaticus* DNA polymerase, *Thermus thermophilus* DNA polymerase, *Escherichia coli* Exo III, and the *Saccharomyces cerevisiae* Rad1/Rad10 complex.

87. (previously presented) The method of Claim 85, wherein said second nuclease is a 5' nuclease derived from a thermostable DNA polymerase altered in amino acid sequence such that it exhibits reduced DNA synthetic activity from that of the wild-type DNA polymerase but retains substantially the same 5' nuclease activity of the wild-type DNA polymerase.

88. (previously presented) The method of Claim 82, wherein said nucleic acid is selected from the group consisting of RNA and DNA.

89. (previously presented) The method of Claim 82, wherein said nucleic acid of step (a) is double stranded.

90. (previously presented) A method for treating nucleic acid, comprising:
- a) providing:
 - i) a purified FEN-1 endonuclease;
 - ii) a source of a target nucleic acid, said target nucleic acid comprising first and second portions, wherein said second portion is downstream of and contiguous to said first portion;
 - iii) a first oligonucleotide comprising a 5' portion complementary to said first portion of said target nucleic acid; and
 - iv) a second oligonucleotide comprising a 3' portion and a 5' portion, wherein said 5' portion is complementary to said second portion of said target nucleic acid;
 - b) generating a cleavage structure wherein at least said 5' portion of said first oligonucleotide is annealed to said first portion of said target nucleic acid and wherein at least said 5' portion of said second oligonucleotide is annealed to said second portion of said target nucleic acid; and
 - c) exposing said cleavage structure to said FEN-1 endonuclease to generate a cleavage product.
91. (previously presented) The method of Claim 90, further comprising the step of d) detecting said cleavage product.
92. (previously presented) The method of Claim 90, further comprising the step of generating a second cleavage structure, said second cleavage structure comprising said cleavage product hybridized to a third oligonucleotide.
93. (previously presented) The method of Claim 92, further comprising the step of exposing said second cleavage structure to said FEN-1 endonuclease to generate a second cleavage product.
94. (previously presented) The method of Claim 93, further comprising the step of detecting said second cleavage product.
95. (previously presented) The method of Claim 90, wherein said FEN-1 endonuclease comprises an archael FEN-1 endonuclease.

96. (previously presented) The method of Claim 95, wherein said archael FEN-1 endonuclease comprises an *Archaeoglobus fulgidus* FEN-1 endonuclease.
97. (previously presented) The method of Claim 96, wherein said *Archaeoglobus fulgidus* FEN-1 endonuclease comprises SEQ ID NO:179.
98. (previously presented) The method of Claim 90, wherein said FEN-1 endonuclease comprises *Pyrococcus woesei* FEN-1 endonuclease.
99. (previously presented) The method of Claim 90, wherein said FEN-1 endonuclease comprises *Methanococcus jannaschii* FEN-1 endonuclease.
100. (previously presented) The method of Claim 90, wherein said FEN-1 endonuclease comprises *Methanobacterium thermoautotrophicum* FEN-1 endonuclease.
101. (previously presented) The method of Claim 90, wherein said FEN-1 endonuclease comprises *Pyrococcus furiosus* FEN-1 endonuclease.
102. (previously presented) The method of Claim 90, wherein said FEN-1 endonuclease comprises a chimerical FEN-1 endonuclease.
103. (previously presented) The method of Claim 102, wherein said chimerical FEN-1 endonuclease comprises at least a portion of an archael FEN-1 endonuclease.
104. (previously presented) The method of Claim 103, wherein said archael FEN-1 endonuclease comprises a *Archaeoglobus fulgidus* FEN-1 endonuclease.
105. (previously presented) The method of Claim 102, wherein said chimerical FEN-1 endonuclease comprises at least a portion of a endonuclease selected from the group consisting of *Pyrococcus woesei* FEN-1 endonuclease, *Methanococcus jannaschii* FEN-1 endonuclease, and *Methanobacterium thermoautotrophicum* FEN-1 endonuclease.
106. (previously presented) The method of Claim 90, wherein said 3' portion of said second oligonucleotide comprises a 3' terminal nucleotide not complementary to said target nucleic acid.

107. (previously presented) The method of Claim 90, wherein said 3' portion of said second oligonucleotide consists of a single nucleotide not complementary to said target nucleic acid.

108. (previously presented) The method of Claim 90, wherein at least one of said first oligonucleotide and said second oligonucleotide is attached to a solid support.